Generalized Isotonic Regression

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Abstract

We present a new computational and statistical approach for fitting isotonic models under convex differentiable loss functions through recursive partitioning. Models along the partitioning path are also isotonic and can be viewed as regularized solutions to the problem. Our approach generalizes and subsumes the well-known work of Barlow and Brunk (1972) on fitting isotonic regressions subject to specially structured loss functions, and expands the range of loss functions that can be used (for example, adding Huber's loss for robust regression). This is accomplished through an algorithmic adjustment to the recursive partitioning approach recently developed for solving large scale l_2 -loss isotonic regression problems (Spouge et al. 2003, Luss et al. 2011). We prove that the new algorithm solves the generalized problem while maintaining the favorable computational and statistical properties of the l_2 algorithm. The results are demonstrated on both real and synthetic data in two settings: fitting count data using negative Poisson log-likelihood loss, and fitting robust isotonic regressions using Huber's loss.

Keywords: isotonic regression, nonparametric regression, regularization path, convex optimization

1 Introduction

In this paper, we generalize recently developed algorithms for solving large-scale isotonic regressions with l_2 loss function (Spouge et al. 2003, Luss et al. 2012) in order to handle a more general class of loss functions. These generalizations allow for fitting isotonic regressions with useful loss functions such as Huber's loss, which was previously impractical for large problems using generic convex optimization solvers. For example, isotonic regression with Huber's loss can be solved

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with generic quadratic programming solvers that suffer due to the large number of constraints in our problems, whereas the algorithm we introduce takes advantage of the structured constraints and is more efficient by orders of magnitude. Isotonic regression is a nonparametric approach for building models whose fits are monotone in their covariates. Such assumptions are natural to applications in biology (Obozinski et al. 2008), ranking (Zheng et al. 2008), medicine (Schell & Singh 1997), statistics (Barlow & Brunk 1972) and psychology (Kruskal 1964). Assume n data observations $(x_1, y_1), ..., (x_n, y_n)$ and a partial order \leq , e.g., the standard Euclidean one where $x_1 \leq x_2$ if and only if $x_{1j} \leq x_{2j}$ coordinate-wise. We index the set of isotonicity constraints implied by the partial order by $\mathcal{I} = \{(i,j) : x_i \leq x_j\}$. Classic isotonic regression considers the l_2 loss function and solves

$$\min\left\{\sum_{i=1}^{n} (\hat{y}_i - y_i)^2 : \hat{y}_i \le \hat{y}_j \quad \forall (i,j) \in \mathcal{I}\right\}$$
(1)

in $\hat{y} \in \mathbf{R}^n$. Throughout the paper we denote by $m = |\mathcal{I}|$ the number of isotonic constraints and d the dimension of data, i.e., $x_i \in \mathbf{R}^d$.

While the assumption of isotonicity is often natural, isotonic regression has not been extensively applied in "modern" applications for two main reasons. As the number of observations n, the data dimensionality d, and the number of isotonicity constraints m increase, problem (1) suffers from computational as well statistical (i.e., overfitting) difficulties. These are reviewed in Luss et al. (2012), where it is argued that the computational difficulties can be overcome using modern algorithms, while overfitting can be addressed by regularizing the problem in (1), i.e., fitting "less complex" isotonic models than the optimal solution of (1). The Isotonic Recursive Partitioning (IRP) algorithm proposed in Spouge et al. (2003) and Luss et al. (2012) (following previous related work by Maxwell & Muckstadt (1985) and Roundy (1986), among others) can easily solve problems with tens of thousands of observations, and is based on recursive partitioning of the covariate space and constructing isotonic models of increasing complexity, thus generating a regu-

larization path of isotonic models in the sense that isotonic models along the path are regularized by the number of partitions made.

In this paper, we focus on a more general form of isotonic regression that minimizes a convex loss function subject to the isotonicity constraints, i.e., we solve

$$\min \{ f(\hat{y}) : \hat{y}_i \le \hat{y}_j \quad \forall (i, j) \in \mathcal{I} \}$$
 (2)

where $f: \mathbf{R}^n \to \mathbf{R}$ is a separable function such that

$$f(\hat{y}) = \sum_{i=1}^{n} f_i(\hat{y}_i), \tag{3}$$

and $f_i: \mathbf{R} \to \mathbf{R}$ is differentiable and convex for all $i=1,\ldots,n$. Typically, $f_i(\hat{y}_i)=g(\hat{y}_i,y_i)$ measures the fit of \hat{y}_i to the observed response. Table 1 provides several examples for the functions $f_i(\cdot)$ that define $f(\cdot)$ above.

p-norm loss, 1	$(\hat{y}_i - y_i)^p$
δ -Huber loss	$ (\hat{y}_i - y_i)^2/2$ for $ \hat{y}_i - y_i < \delta$ and $\delta(\hat{y}_i - y_i - \delta/2)$ otherwise
Negative Poisson log-likelihood	$\hat{y}_i - y_i \ln \left(\hat{y}_i \right)$
Negative Bernoulli log-likelihood	$-y_i \ln(\hat{y}_i) - (1 - y_i) \ln(1 - \hat{y}_i)$

Table 1: Examples of loss functions solvable by Algorithm 1

The notion of generalized isotonic regression is not new. Barlow & Brunk (1972) defined a generalized loss function as

$$f(\hat{y}) = \sum_{i=1}^{n} \Phi(\hat{y}_i) - \hat{y}_i y_i \tag{4}$$

for proper convex $\Phi : \mathbf{R} \to \mathbf{R}$ which was minimized subject to the same isotonicity constraints as in (2). They showed that this generalized isotonic regression problem can be solved equivalently as an instance of the l_2 isotonic regression (1). This implies that any large-scale algorithm for problem (1) can be used to solve isotonic regressions with objectives of the form (4). Generalized objectives

for large-scale Poisson and Bernoulli regressions as given in Table 1 can be solved in this manner, however the p-norm and Huber loss functions cannot. This relationship will be further formalized in Section 2.4. Generalized isotonic regression (using separable loss functions) in d=1 dimension was also considered in Best et al. (2000) and Ahuja & Orlin (2001) using extensions of the pooled adjacent violators algorithm (PAVA). Neither assumes differentiability as done here, and hence both are amenable to a broader class of loss functions, albeit only in one dimension. Hochbaum & Queyranne (2003) offer a very efficient and related algorithm for problem (2) where the fits are restricted to being integer (called the convex cost closure problem). Their algorithm can be extended to the continuous case in the sense of determining an ϵ -accurate solution by solving the integer problem on an ϵ -grid. Depending on the required level of accuracy, their approach can be computationally competitive with our algorithm as described below for finding the solution of (2), however it lacks the natural statistical interpretation as a regularization path which our approach affords. The method of Hochbaum & Queyranne (2003) is discussed in more detail in Section 2.4.

The main contribution of this paper is a generalization of IRP that can be used to solve large-scale multivariate generalized isotonic regressions of the form (2). Our generalization extends the methods to any convex differentiable loss function, including those mentioned in Table 1, and we term it generalized isotonic recursive partitioning (GIRP). As with IRP, the partitioning algorithm here addresses both of the main difficulties with isotonic regression discussed above. Firstly, it provides a sequence of isotonic models of increasing complexity, converging to the globally optimal generalized isotonic solution. Early stopping along this "regularization path" is a useful approach to overcome overfitting concerns of the globally optimal solution; less complex isotonic models along the path often predict more accurately than the final overfit model. Secondly, it is computationally efficient; the partitioning algorithm is an iterative scheme in which each iteration partitions a group of observations by solving a structured linear program for which very efficient algorithms exist.

It should be emphasized that, while the algorithmic modification from IRP to GIRP is quite

minor and the proofs for GIRP properties are closely related to proofs of same results for IRP, the generalization has important practical implications because it significantly expands the range of applications for isotonic modeling, as our examples below illustrate.

The paper continues as follows. Section 2 describes the known results for l_2 isotonic regression and generalizes them to the class of convex loss functions described by (3). The generalized algorithm is described, and the relationship to Barlow & Brunk (1972) is formalized. Section 3 applies the results with Poisson log-likelihood and Huber's loss functions to synthetic and real data sets. A Matlab-based software package implementing our results is available at www.tau.ac.il/~saharon/files/GIRPv1.zip. We first define terminology to be used throughout the remainder of the paper.

1.1 Definitions

Let $V = \{x_1, \dots, x_n\}$ be the covariate vectors for n training points where $x_i \in \mathbb{R}^d$ and denote $y_i \in \mathbb{R}$ as the i^{th} observed response. We will refer to a general subset of points $A \subseteq V$ with no holes (i.e., $x \preceq y \preceq z$ and $x, z \in A \Rightarrow y \in A$) as a *group*. Throughout the paper, we will use the shorthand $i \in A = \{i : x_i \in A\}$. Denote by |A| the cardinality of group A. The *weight* of a group A is denoted by

$$w_A = \underset{z \in \mathbf{R}}{\operatorname{argmin}} \sum_{i \in A} f_i(z). \tag{5}$$

For two groups A and B, we denote $A \leq B$ if $\exists x \in A, y \in B$ such that $x \leq y$ and $\nexists x \in A, y \in B$ such that $y \prec x$ (i.e., there is at least one comparable pair of points that satisfy the direction of isotonicity). A set of groups \mathcal{V} is called isotonic if $A \leq B \Rightarrow w_A \leq w_B, \forall A, B \in \mathcal{V}$. A subset \mathcal{L} (\mathcal{U}) of A is a lower set (upper set) of A if $x \in A, y \in \mathcal{L}, x \prec y \Rightarrow x \in \mathcal{L}$ ($x \in \mathcal{U}, y \in A, x \prec y \Rightarrow x \in \mathcal{L}$). A group $B \subseteq A$ is defined as a block of group A if $w_{\mathcal{U} \cap B} \leq w_B$ for each upper set \mathcal{U} of A such that $\mathcal{U} \cap B \neq \{\}$ (or equivalently if $w_{\mathcal{L} \cap B} \geq w_B$ for each lower set \mathcal{L} of A such that $\mathcal{L} \cap B \neq \{\}$). We denote the optimal solution for minimizing f(y) in the variable y by y^* , i.e.,

 $y^* = \operatorname{argmin} f(y)$. The notation $(\partial f(\hat{y})/\partial \hat{y})|_y$ denotes the derivative of a function f with respect to the variable \hat{y} evaluated at the point y.

2 Generalized Isotonic Recursive Partitioning with Convex Loss Functions

In this section, we generalize the results for l_2 isotonic partitioning resulting in an algorithm termed Generalized Isotonic Recursive Partitioning (GIRP), and derive useful properties of this generalization. The solution at each iteration, as in IRP, is defined by groups that are proven to be the union of blocks in the optimal solution. Section 2.1 first gives an overview of the IRP algorithm. Section 2.2 then details the partitioning step for the generalized case and derives the resulting GIRP algorithm. When $f_i(\hat{y}_i) = (\hat{y}_i - y_i)^2$, that is, $f(\cdot)$ is the l_2 loss function, all results in this section replicate those of IRP and w_A becomes the average of the observations in group A. Section 2.3 proves convergence of the partitioning algorithm to the global optimal solution of (2) and shows that the solution at each iteration of the algorithm is isotonic, i.e., the iterations provide a regularization path of isotonic solutions.

2.1 Isotonic Recursive Partitioning with the l_2 Loss Function

We here briefly review the ideas of the IRP algorithm of Spouge et al. (2003) and Luss et al. (2012). The optimal solution to the l_2 isotonic regression problem (1) is known to be defined by a partitioning of the observations y_i into blocks in which $\hat{y}_i = \hat{y}_j$ if observations y_i and y_j are in the same block. Indeed, this structure can be seen through the optimality conditions (i.e., Karush-Kuhn-Tucker (KKT) conditions, see Boyd & Vandenberghe (2004)) for problem (1). The optimal solution to (1), denoted by \hat{y}^* , satisfies these conditions, which are given by

(a)
$$2(\hat{y}_i^* - y_i) = \sum_{j:(j,i)\in\mathcal{I}} \lambda_{ji}^* - \sum_{j:(i,j)\in\mathcal{I}} \lambda_{ij}^* \quad \forall i \in V$$

(b)
$$\hat{y}_i^* \leq \hat{y}_i^* \quad \forall (i,j) \in \mathcal{I}$$

(c)
$$\lambda_{ij}^* \ge 0 \quad \forall (i,j) \in \mathcal{I}$$

(d)
$$\lambda_{ij}^*(\hat{y}_i^* - \hat{y}_j^*) = 0 \quad \forall (i,j) \in \mathcal{I},$$

where λ_{ij}^* is the optimal dual variable associated with isotonicity constraint $\hat{y}_i \leq \hat{y}_j$. Convexity of the l_2 loss function implies that any solution satisfying conditions (a)-(d) is a globally optimal solution. From condition (d), $\lambda_{ij}^* > 0 \Rightarrow \hat{y}_i^* = \hat{y}_j^*$, i.e., the optimal solution is made of blocks V where $\lambda_{ij}^* > 0$ for all $i, j \in V$, which implies that all observations within a block V are fit to the same value. Furthermore, when restricting all fits within a block V to be equivalent, the isotonic regression problem over block V is the following unconstrained quadratic program

$$\min_{\hat{y} \in \mathbf{R}} \left\{ \sum_{i \in V} (\hat{y} - y_i)^2 \right\} \tag{6}$$

which is trivially solved at $\hat{y}^* = \overline{y}_V$ where \overline{y}_V denotes the average of all observations in block V. Condition (b) implies that these averages must satisfy isotonicity, i.e., if $V^- \leq V^+$ are isotonic blocks then $\overline{y}_{V^-} \leq \overline{y}_{V^+}$. Thus, the structure of the optimal solution is a partitioning of the set $\{1,\ldots,n\}$ into some (unknown number) K blocks $\{V_1,\ldots,V_K\}$ where $\hat{y}_i^* = \overline{y}_{V_k}$ for all $i \in V_k$ and $\hat{y}_i^* \leq \hat{y}_j^*$ for all $(i,j) \in \mathcal{I}$.

Many such feasible partitions exist that are not optimal (e.g., set all fits to the average of all observations). Condition (a) above must also be satisfied and gives the motivation for how to partition the set of observations in a manner that leads to the optimal partitioning. The partitioning scheme is detailed for the general case of convex loss functions in the next subsection. Here, we only give a general idea of how IRP works.

IRP starts with the entire dataset as one group V and iteratively splits it into an increasing number of groups, until the optimal solution of (1) is reached. At each iteration, the algorithm chooses a sub-optimal group and partitions it into two groups by solving a specially structured

linear program, detailed in the next subsection, that is amenable to very efficient algorithms. If the partition puts all observations into one group, it can be shown that the group is a block, i.e., optimal. Otherwise, the fits in the two resulting groups are recalculated as their averages (via (6) above), while the rest of the groups and their fits remain at their values in the previous iteration. IRP is thus an iterative scheme that splits a group at each iteration and never merges two groups back together; therefore, IRP is referred to as a no-regret partitioning algorithm.

Two important theorems are proven (Luss et al. 2012) with respect to IRP. The first states that the new solution obtained after each partitioning step still satisfies isotonicity. After iteration K, there are K+1 groups, V_1, \ldots, V_{K+1} , in the partitioning with \hat{y}_i fit to \overline{y}_{V_k} for all $i \in V_k$ with $k \in \{1, \ldots, K+1\}$. The theorem thus says, that at each iteration K, the fits from the partitioning V_1, \ldots, V_{K+1} provide a potential isotonic prediction model. The second theorem shows that the IRP scheme terminates at the globally optimal partitioning. Hence, IRP produces a path of increasingly complex (since each iteration adds a partition) isotonic solutions, terminating in the optimal solution of (1). These theorems are made possible because of the particular splitting criterion used in the IRP algorithm, which is amenable to efficient calculation as mentioned above. The generalized version of the splitting criterion and the resulting algorithm are discussed next.

2.2 The partitioning algorithm

As with IRP, we solve a sequence of subproblems in order to solve the generalized isotonic regression problem (2); each subproblem divides a group of observations into two groups at each iteration. An important property of IRP with the l_2 loss function is that observations separated at one iteration remain separated at all future iterations. The same property applies here and implies that the total number of iterations is bounded by the number of observations n.

The partitioning algorithm is motivated by the optimality conditions for the generalized isotonic regression problem (2). The optimal solution to (2), denoted by \hat{y}^* , are identical to conditions (a)-(d) in Section 2.1 above, with the exception that condition (a) now has the generalized form

(a)
$$\frac{\partial f_i(\hat{y}_i)}{\partial \hat{y}_i}\Big|_{\hat{y}_i^*} = \sum_{j:(j,i)\in\mathcal{I}} \lambda_{ji}^* - \sum_{j:(i,j)\in\mathcal{I}} \lambda_{ij}^* \quad \forall i \in V$$

where again λ_{ij}^* is the optimal dual variable associated with isotonicity constraint $\hat{y_i} \leq \hat{y_j}$. Convexity of the loss function again implies that any solution satisfying the optimality conditions is a globally optimal solution. The structure of the optimal solution as a partitioning of isotonic blocks can be seen from the KKT conditions as described in Section 2.1. Within each block, the fit to each observation for the general case is taken to be the weight of the observations in the block as defined by (5). Isotonicity of the two blocks V^- and V^+ , i.e., $V^- \leq V^+$, means that $w_{V^-} \leq w_{V^+}$. From condition (a), summing over all observations in a block V, i.e., optimal group, gives

$$\sum_{i \in V} \frac{\partial f_i(\hat{y}_i)}{\partial \hat{y}_i} \bigg|_{\hat{y}_i^*} = 0. \tag{7}$$

Derivation of the partitioning step is as follows. Consider a group V where $\hat{y}_i^* = w_V$ for all $i \in V$. If V is an optimal group, it is a block and must satisfy (7). If it is not optimal, however, we can find a partitioning of V into two isotonic groups V^+ and V^- such that

$$\sum_{i \in V^{+}} \frac{\partial f_{i}(\hat{y}_{i})}{\partial \hat{y}_{i}} \bigg|_{w_{V}} - \sum_{i \in V^{-}} \frac{\partial f_{i}(\hat{y}_{i})}{\partial \hat{y}_{i}} \bigg|_{w_{V}} < 0.$$
(8)

The first summation over $i \in V^+$ is the change in the objective value of problem (2) due to an increase in the fits of observations in V^+ . The second summation over $i \in V^-$ is the change in the objective value due to a decrease in the fits of observations in V^- . Such a partition thus means that increasing the fits in V^+ to be greater than w_V while decreasing the fits in V^- to be less than w_V (which by definition maintains isotonicity of the fits) will cause an overall decrease in the objective value to problem (2). Fits that decrease the overall objective value can be achieved by fitting the observations in V^+ and V^- to their respective weights w_{V^+} and w_{V^-} . Hence, we search for an isotonic partitioning of V into V^+ and V^- that minimizes the lefthand term in (8).

Denote by
$$C_V = \{(V^-, V^+) : V^-, V^+ \subseteq V, V^- \cup V^+ = V, V^- \cap V^+ = \{\}, \not\exists x \in V^-, y \in V^+ \cap V^+ = \{\}, \forall x \in V^-, y \in V^+ \cap V^+ = \{\}, \forall x \in V^-, y \in V^+ \cap V^+ = \{\}, \forall x \in V^-, y \in V^+ \cap V^+ = \{\}, \forall x \in V^-, y \in V^+ \cap V^+ = \{\}, \forall x \in V^-, y \in V^+ \cap V^+ = \{\}, \forall x \in V^-, y \in V^+ \cap V^+ = \{\}, \forall x \in V^-, y \in V^+ \cap V^+ = \{\}, \forall x \in V^-, y \in V^+ \cap V^+ = \{\}, \forall x \in V^-, y \in V^+ \cap V^+ = \{\}, \forall x \in V^-, y \in V^+ \cap V^+ = \{\}, \forall x \in V^-, y \in V^+ \cap V^+ = V^+ \cap V^+ = \{\}, \forall x \in V^-, y \in V^+ \cap V^+ = V^+ \cap V^+ = V^+ \cap V^+ = V^+ \cap V^+ = V^+ \cap V^+ \cap V^+ \cap V^+ = V^+ \cap V^+ \cap V^+ = V^+ \cap V^+ \cap V^+ \cap V^+ = V^+ \cap V^+ \cap V^+ \cap V^+ \cap V^+ = V^+ \cap V^+ \cap V^+ \cap V^+ \cap V^+ \cap V^+ = V^+ \cap V^+$$

 V^+ s.t. $y \leq x$ the set of all feasible (i.e., isotonic) partitions defined by observations in V. Partitioning is referred to as making a cut through the variable space (hence the optimal partition is made by an *optimal cut*). The optimal cut is determined as the partition that solves the problem

$$\min_{(V^-,V^+)\in\mathcal{C}_V} \left\{ \sum_{i\in V^+} \frac{\partial f_i(\hat{y}_i)}{\partial \hat{y}_i} \bigg|_{w_V} - \sum_{i\in V^-} \frac{\partial f_i(\hat{y}_i)}{\partial \hat{y}_i} \bigg|_{w_V} \right\} \tag{9}$$

where $V^-(V^+)$ is the group on the lower (upper) side of the edges of the cut. The optimal cut problem (9) can be expressed as the binary program

$$\min \left\{ \sum_{i \in V} x_i \frac{\partial f_i(\hat{y}_i)}{\partial \hat{y}_i} \right|_{w_V} : x_i \le x_j \quad \forall (i, j) \in \mathcal{I}, x_i \in \{-1, +1\} \quad \forall i \in V \}.$$
 (10)

It is well-known (Murty 1983) that the continuous relaxation to this binary program (i.e., replacing the constraints $x_i \in \{-1, +1\}$ by $-1 \le x_i \le 1$ for all $i \in V$) is solved on the boundary of the feasible region with $x_i^* \in \{-1, +1\}$ for all $i = 1 \dots n$. Thus the optimal cut problem (9) is equivalent to solving the linear program

$$\min \left\{ z^T x : x_i \le x_j \quad \forall (i,j) \in \mathcal{I}, -1 \le x_i \le 1 \quad \forall i \in V \right\}$$
 (11)

where $z_i = (\partial f_i(\hat{y}_i)/\partial \hat{y}_i)|_{w_V}$. Problem (11) with $z_i = 2(\overline{y}_V - y_i)$ gives the linear program used to make partitions in IRP with l_2 loss function as described above in Section 2.1. As seen by property (8), a property of this optimal cut for generalized isotonic regression is that the sum of loss functions with $x_i = +1$ ($x_i = -1$) can be decreased by increasing (decreasing) the corresponding fits. That is, by increased (decreasing) w_v for observations i with $x_i = +1$ ($x_i = -1$), the total change in loss is decreased, i.e.,

$$\sum_{\{i:x_i=+1\}} \frac{\partial f_i(\hat{y}_i)}{\partial \hat{y}_i} \bigg|_{w_V} \le 0 \quad \text{and} \quad \sum_{\{i:x_i=-1\}} \frac{\partial f_i(\hat{y}_i)}{\partial \hat{y}_i} \bigg|_{w_V} \ge 0.$$
 (12)

This group-wise partitioning operation is the basis for our algorithm which is detailed in Algorithm 1. The algorithm differs from IRP only in Step 6 (they are obviously identical when $f(\cdot)$ is the l_2 loss). Initially, all observations are in one group. Each iteration splits a group optimally by solving subproblem (11). A list \mathcal{C} of potential optimal cuts for each group generated thus far is maintained, and, at each iteration, the cut among them with the smallest (most negative) objective value is performed. Partitioning of a group ends when the solution to (11) is trivial (i.e., no split is found because the group is a block). The algorithm stops when no further groups can be partitioned.

Algorithm 1 Generalized Isotonic Recursive Partitioning

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Require: Observations (x_1, y_1), \ldots, (x_n, y_n) and partial order \mathcal{I}.
Require: k = 0, \mathcal{A} = \{\{x_1, \dots, x_n\}\}, \mathcal{C} = \{(0, \{x_1, \dots, x_n\}, \{\})\}, \mathcal{B} = \{\}, M_0 = (\mathcal{A}, w_{\mathcal{A}}).
  1: while A \neq \{\} do
          Let (val, w^-, w^+) \in \mathcal{C} be the potential partition with largest val.
          Update \mathcal{A} = (\mathcal{A} \setminus (w^- \cup w^+)) \cup \{w^-, w^+\}, \mathcal{C} = \mathcal{C} \setminus (val, w^-, w^+).
          M_k = (\mathcal{A}, \overline{y}_{\Delta}).
 4:
          for all v \in \{w^-, w^+\} \setminus \{\} do
 5:
             Set c_i = \frac{\partial f_i(\hat{y}_i)}{\partial \hat{y}_i}\big|_{w_v} \ \forall i \in v \ \text{where} \ w_v \ \text{is the weight (5) of the observations in } v.
Solve LP (11) with input z and get z^* = \operatorname{argmin} \operatorname{LP}(11).
 6:
 7:
              if z_1^* = \ldots = z_n^* (group is optimally divided) then
 8:
                  Update A = A \setminus v and B = B \cup \{v\}.
 9:
10:
                 Let v^- = \{x_i : z_i^* = -1\}, v^+ = \{x_i : z_i^* = +1\}.
11:
                 Update \mathcal{C} = \mathcal{C} \cup \{(c^T z^*, v^-, v^+)\}
12:
              end if
13:
          end for
14:
          k=k+1.
15:
16: end while
17: return \mathcal{M}, a sequence of isotonic models, where M_k contains the k^{th} iteration's partitioning
      of observations and corresponding group weights.
```

Each iteration k of Algorithm 1 produces a model M_k by fitting each group in M_k to its weight. For a set of groups $\mathcal{V} = \{V_1, \dots, V_k\}$, denote $w_{\mathcal{V}} = \{w_{V_1}, \dots, w_{V_k}\}$. Then model $M_k = (\mathcal{V}, w_{\mathcal{V}})$ contains the partitioning \mathcal{V} as well as a fit to each of the observations, which is the weight, as defined by (5), of the group it belongs to in the partition.

2.3 Properties of the partitioning algorithm

So far, we have detailed the partitioning algorithm which is based on iteratively solving problem (11), but we have not yet shown that partitioning according to this particular scheme, i.e., solving problem (11), optimally solves the generalized isotonic regression problem. Theorem 1 next states the main result that implies Algorithm 1 is a no-regret partitioning algorithm for (2) (no-regret in the same sense as described for IRP in Section 2.1). In the case of l_2 isotonic regression, this result is already known (Maxwell & Muckstadt 1985, Spouge et al. 2003, Luss et al. 2012). This theorem leads to our convergence result. The proof requires straightforward changes to the proof in Luss et al. (2012) based on the definition of convexity, the new algorithm cut in (11), and its properties (12); the proof is thus left to the Appendix.

Theorem 1 Assume group V is the union of blocks from the optimal solution to problem (2). Then a cut made by solving (11) at a particular iteration does not cut through any block in the global optimal solution.

The case of multiple observations at the same coordinates can be disregarded. Let \mathcal{V} denote a set of groups where each group in \mathcal{V} contains observations with the same coordinates, i.e $V \in \mathcal{V}$ denotes the indices of multiple observations and |V|=1 means that V is a single observation. Then, we define $g_V(\hat{y}_V)=\sum_{i\in V}f_i(\hat{y}_V)$ where $\hat{y}_V\in\mathbf{R}$ and modify $f(\cdot)$ in the generalized isotonic regression problem (2) to be $f(\hat{y})=\sum_{V\in\mathcal{V}}g_V(\hat{y}_V)$ where $\hat{y}\in\mathbf{R}^{|\mathcal{V}|}$ and each function $g_V(\cdot)$ satisfies the necessary properties for applying GIRP.

Since Algorithm 1 starts with the union of all blocks for the first partition, we can conclude from this theorem that Algorithm 1 never cuts a block when generating partitions. From the derivation of the partitioning problem, it is clear that if an isotonic partition can be made, it will be made; that is, the algorithm will not stop early. Convergence of Algorithm 1 to the global isotonic solution with no regret then follows by repeatedly applying Theorem 1 until all blocks of the optimal solution are identified. The next theorem states that Algorithm 1 provides isotonic solutions at each

iteration. This result implies that the path of solutions generated by Algorithm 1 can be regarded as a regularization path for the generalized isotonic regression problem (2). Proof of this theorem is again held until the Appendix for the same reasons given above.

Theorem 2 *Model* M_k *generated after iteration* k *of Algorithm 1 is in the class of isotonic models.*

Complexity analysis of Algorithm 1 depends on the number of observations n and isotonic constraints m, and the complexity of solving linear program (11). Firstly, we assume that computing the weight of a group V via (5) requires computationally less effort than solving problem (11) (in practice these problems are one-dimensional convex minimization problems that are easily solved with a binary search). In short, linear program (11) is dual to a linear maximum flow network problem (Ahuja et al. 1993), which is a well-studied problem. It can be solved in $O(mn \log n)$ (Sleator & Tarjan 1983) or $O(n^3)$ (Galil & Naamad 1980) in the general case that we consider; special cases such as n=2 or where the observations lie on a grid can be computed even faster (Spouge et al. 2003). Choice of algorithm depends on m which is $O(n^2)$ in the worst case. Given GIRP requires at most n iterations, this leads to worst case complexities of $O(mn^2 \log n)$ or $O(n^4)$. A recent problem reduction by Stout (2010) can be used to obtain an equivalent representation of the desired problem with d-dimensional data and $O(n \log^{d-1} n)$ constraints and observations, which can be useful when m is large. Finally, Luss et al. (2012) show that IRP performs in $O(Cn^3)$ in practice, where C is a function of the fraction of observations on each of the cut at each iteration. The same result applies here.

2.4 Relations to Other Generalized Isotonic Regressions

We here formalize the relationship between GIRP and the work of Barlow & Brunk (1972), which was hinted at in Luss et al. (2012) and mentioned in the introduction above. The generalized

isotonic regression problem of Barlow & Brunk (1972) is of the form

$$\min\left\{\sum_{i=1}^{n} \Phi(\hat{y}_i) - \hat{y}_i y_i : \hat{y}_i \le \hat{y}_j \quad \forall (i,j) \in \mathcal{I}\right\}$$
(13)

in $\hat{y} \in \mathbf{R}^n$ where we have left out weights for simplicity. While they allow $\Phi : \mathbf{R} \to \mathbf{R}$ to be nondifferentiable, we consider here only the differentiable case and denote $\phi(\cdot)$ as the derivative of $\Phi(\cdot)$. Let \hat{z}^* be the solution of (1) (i.e., l_2 isotonic regression) with given observations y_i . Theorem 3.1 of Barlow & Brunk (1972) claims that the solution to (13) can be obtained as

$$\hat{y}_i^* = \phi^{-1}(\hat{z}_i^*) \tag{14}$$

where $\phi^{-1}(\cdot)$ is defined by $\phi^{-1}(\phi(x)) = x$ for all $x \in \mathbf{R}$. Thus, any objective of the form (13) can be solved by computing the l_2 isotonic regression on input observations and then transforming the solution using (14). In this manner, IRP can be used to solved the somewhat limited class of generalized isotonic regression problems defined in Barlow & Brunk (1972) (note that without requiring $\Phi(\cdot)$ proper, their theory would apply to any convex loss function). However, it is also clear that Algorithm 1 provides the tools for solving more general isotonic regression problems than (13), e.g., as in the case for the p-norm or Huber's loss function.

The same transformation from Barlow & Brunk (1972) can be used to derive an isotonic regularization path for generalized isotonic regression problems with the structure of (13). Indeed, this can be shown using the above framework for our general isotonic regression problem (2), and is formalized in Proposition 3.

Proposition 3 *Problem (13) can be solved either by*

- 1. Applying IRP to the observation data y to obtain \hat{z}^* and tranforming using (14),
- 2. Applying GIRP directly to (13).

Furthermore, both algorithms are equivalent when applied to (13) in the sense that the regulariza-

tion path of partitions for each algorithm are equivalent.

Proof. IRP can be used to solve the l_2 isotonic regression problem to obtain \hat{z}^* . Application of Theorem 3.1 of Barlow & Brunk (1972) gives the solution to (13) via (14). In order to apply GIRP, let $f_i(\hat{y}_i) = \Phi(\hat{y}_i) - \hat{y}_i y_i$ and $(\partial f_i(\hat{y}_i)/\partial \hat{y}_i)|_{\hat{y}_i} = \phi(\hat{y}_i) - y_i$ where $\phi(\cdot)$ denotes the derivative of $\Phi(\cdot)$. Then w_V as defined by (5) satisfies $\sum_i (\phi(w_V) - y_i) = 0$ giving $w_V = \phi^{-1}(\overline{y}_V)$ where \overline{y}_V denotes the mean observation over group V and $\phi^{-1}(\cdot)$ is defined by $\phi^{-1}(\phi(x)) = x$ for all $x \in \mathbb{R}$. Hence, here $(\partial f_i(\hat{y}_i)/\partial \hat{y}_i)|_{w_V} = \overline{y}_V - y_i$ and the GIRP partitioning problem (11) is equivalent to the corresponding partition problem in Luss et al. (2012).

The only difference between IRP and GIRP for solving problem (13) is that GIRP fits observations to the transformed isotonic regression fits $w_V = \phi^{-1}(\overline{y}_V)$ along the path, while IRP fits observations to the mean of the group's observations and the transformation is done on the final optimal partitioning. It is easy to see that the transformation (14) can be applied to each iteration of IRP along the path in order to obtain an equivalent path to that of GIRP.

This connection also relates the algorithm of Maxwell & Muckstadt (1985), which solves problem (13) with $\Phi(\hat{y}_i) = 1/\hat{y}_i$, to IRP. Due to the analysis here, these algorithms are actually equivalent. Both the problem of Maxwell & Muckstadt (1985) and l_2 isotonic regression are specific instances of the more general problem (2) solved in this paper. It should be noted that Maxwell & Muckstadt (1985) did not make use of, or even recognize, the regularization path which plays a significant role for isotonic regression in dimension d > 1.

Lastly, Hochbaum & Queyranne (2003) offer another partitioning algorithm for problem (2) with additional integer constraints. GIRP, in the continuous case, solves cut problem (9) because we know the fit within optimal groups (i.e., the weight of the group). Rather, in the integer case, the cut problem (9) is solved instead with the derivatives evaluated at some α taken as the median of an interval in which the optimal fits lie. A theorem states that this partition problem divides the group into two groups V^- and V^+ , where optimal fits to observations in V^- are less than α and

optimal fits to observations in V^+ are greater than α . The problem is thus stated as determining a sequence $\alpha_1, \ldots, \alpha_l$ such that observations with optimal fits in the interval $(\alpha_i, \alpha_{i+1}]$ have the same optimal fit. Given a criterion for determining when α is a breakpoint in this sequence, their algorithm can do better than a binary search. In fact, they further suggest a method that has a worst-case complexity equivalent to solving three max-flow problems. The complexity comes from using the information in previous max-flow problems to start new max-flow problems. A similar idea could possibly be applied in our continuous case where the search for breakpoints uses the group weight in the cut problem. This highly efficient algorithm does not provide the exact solution to the continuous case, but a regularization path based on the bounds they get when searching for breakpoints can be considered for the integer case, and in turn, for the problem on an ϵ -grid.

2.5 Regularization by recursive partitioning

GIRP obtains the solution to problem (2) by recursively partitioning the covariate space into progressively smaller regions and fitting the best constant in each region, referred to here as the weight which is defined by (5). As such, it is natural to think of the resulting sequence of models created from early stopping as a regularization path of models of increasing complexity, indexed by the number of iterations of the algorithm. Other examples of using early stopping for regularization include training neural networks with back propagation (Caruana et al. 2000) and boosting (Rosset et al. 2004). Extensive experience of the usefulness of regularization in high dimensional fitting (Wahba 1990, Tibshirani 1996, Schölkopf & Smola 2001), and especially in nonparametric models like isotonic regression, suggests that regularization, embodied in this case by early stopping of the algorithm, can lead to reduced overfitting and hence improved predictive performance. As Theorem 2 indicates, when stopping early and fitting the weight to each region, we are guaranteed to obtain a feasible isotonic model.

While GIRP uses early stopping for regularization of the globally optimal isotonic model, we note that regularization commonly refers to learning a model by explicitly constraining the family of models that are considered, and optimizing over this family. Early stopping after the k^{th} iteration of GIRP produces an isotonic model with k cuts obtained through a sequence of local optimization problems. However, this model is not the solution to any global optimization problem. The k^{th} model of GIRP is thus only one potential model with k cuts that satisfies the isotonicity constraints. One might, for example, seek a regularized isotonic model that minimizes loss subject to the isotonicity constraints such that exactly k cuts are made. The k^{th} model in this case has a clear interpretation and more flexibility than the k^{th} GIRP model. While this would certainly be an interesting problem to consider, it is combinatorially difficult and the authors do not know of any efficient methods for solving it.

In Luss et al. (2012), model complexity of l_2 isotonic regression along the IRP path is quantified through the concept of equivalent degrees of freedom (DFs) as defined by Efron (1986) and Hastie et al. (2001). The initial iterations of IRP are shown to perform much more fitting than later iterations, and this phenomenon becomes more pronounced as the dimension d increases. For example, in dimension 6, often 50% of DFs were fitted by the first IRP iteration. Although the model complexity and DF measures of Efron (1986) do not generalize to non- l_2 loss as used in GIRP, the general spirit of this result should persist. Intuitively, because the space of isotonic splits of the entire covariate space probed in the first iteration is much larger than the space of possible isotonic cuts in further iterations, finding the optimal first split corresponds to a significant portion of all fitting.

These two effects — importance of early stopping coupled with the high portion of fitting in earlier iterations — are demonstrated empirically in the experiments of the next section, where the best performing solution along the GIRP path is compared to the optimal solution of problem (2) in terms of predictive performance.

3 Performance evaluation

We here demonstrate usefulness of the partitioning algorithm for generalized loss functions. The contribution of our generalization is specifically illustrated by the use of Huber loss, which proves to be very effective in the case of outliers. We first exhibit the computational performance of GIRP and show that the algorithm can be applied to large-scale problems. We then consider synthetic data sets that demonstrate the impact of regularization and conclude with an example on real data.

3.1 Practical Computational Performance

Solving the multivariate isotonic regression problem with general loss functions such as Huber's loss was previously a computationally difficult problem. For certain loss functions, the isotonic regression problem can be reformulated and solved with off-the-shelf convex optimization solvers. For example, isotonic regression with Huber's loss can be reformulated as a quadratic program by adding many variables to the optimization problem. Simulations with 1000 training points were solved in 2.3 seconds with GIRP versus 135 seconds using Mosek (MOSEK ApS 2011) to solve the quadratic program (averaged over 50 simulations). This simple experiment demonstrates that GIRP, which is specifically designed for isotonic regression problems, is clearly a much more practical tool than using off-the-shelf generic solvers and makes generalized isotonic regression problems amenable to large-scale problems.

Figure 1 (left) illustrates that GIRP can solve large-scale problems with Huber's loss. The i^{th} observation in each simulation is generated as $y_i = (\prod_j x_{ij}) + \mathcal{N}(0, d^2)$ with $x_{ij} \sim \mathcal{U}[0, 2]$, d representing the dimension, and outliers randomly inserted. Results are averages over 50 simulations. Isotonic regression in 8 dimensions with 20,000 training instances is solved in less than one minute. Figure 1 (right) shows the number of partitions that GIRP performs on average for varying dimensions. More training data and higher dimension typically implies more complex isotonic models, resulting in more partitioning problems and more computational time. The com-

putational limitation of training the isotonic model with GIRP is solving the partitioning problem. Luss et al. (2010) further offers a heuristic for solving the partition problem that makes training isotonic regression problems with up to 200,000 training instances easily feasible.

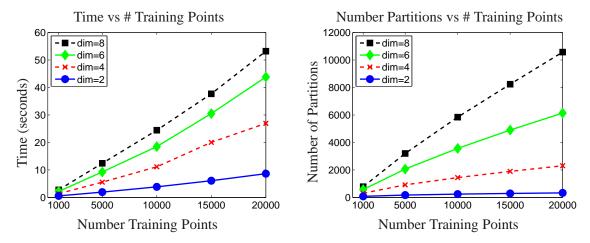


Figure 1: Left: Computational performance of training isotonic models with GIRP on a simulation with Huber's loss for varying number of dimensions and training instances. Right: Complexity of isotonic models as measured by the number of partitions for varying number of dimensions and training instances.

3.2 Simulations

Experiments are run on two different loss functions. In the first experiment, count data is simulated from Poisson distributions where the average number of occurrences is generated by two different isotonic models. Generalized isotonic regression models for the Poisson rate are obtained by minimizing negative Poisson log-likelihood subject to isotonicity constraints. In the second set of experiments, observations are generated by two different isotonic models and .5% of the training observations are multiplied by a large constant to make them outliers. Generalized isotonic regression models are obtained using δ -Huber loss. Note that Poisson isotonic regressions can be handled using IRP due to the theory of Barlow & Brunk (1972), while Huber isotonic regressions require using GIRP.

Our experimental framework is as follows. A training and testing set are independently simulated by a fixed distribution. Training and testing sets have 15000 and 3000 observations, respec-

tively. A model is first generated on the training data. In the case of GIRP, the training data is split into a subtraining set of 12000 observations and a validation set of the remaining 3000 observations. A path of isotonic models is generated by running GIRP on the subtraining data. The validation data is used to select the regularization level (stopping point), and the resulting model is applied to predict the testing data. With respect to parametric regression, e.g., Poisson and Huber regressions, models are trained on the full 15000 observation training set and tested on the 3000 observation testing set. Results are based on averaging fifty simulations.

The first two examples use Poisson negative log-likelihood as the loss function. Data for the two simulations is generated as $x_{ij} \sim \mathcal{U}[0,10]$ and $x_{ij} \sim \mathcal{U}[5,10]$ (the coordinates of x are drawn i.i.d in all our experiments), respectively. The i^{th} observation in each simulation is generated as $y_i \sim \text{Poisson}(\prod_j \sqrt{x_{ij}})$ and $y_i \sim \text{Poisson}(\sum_j x_{ij}^2)$, respectively. The isotonic models are compared to the results of a Poisson regression, and performance here is measured by negative Poisson log-likelihood. The regularized model generated by the minimum loss along the GIRP curve (GIRP Min Poisson) is compared with the final GIRP model (GIRP Final Poisson) and with the Poisson regression model. In practice, one would only consider predictions using the regularized model, but here we want to compare against the unregularized model as well. Table 2 demonstrates that Poisson isotonic regression works well with a reasonable number of variables (2-5 for the first simulation and 2-3 for the second simulation), however is outperformed by the simple Poisson regression with more than 5 variables. In comparing the regularized GIRP model with the final GIRP model, there is no statistical difference in this example. The next simulation clearly exemplifies the effect of regularization, in addition to the use of generalized isotonic regression.

The second two examples use δ -Huber loss as the loss function for generating models. Data for the two simulations is generated as $x_{ij} \sim \mathcal{U}[0,3]$ and $x_{ij} \sim \mathcal{U}[0,5]$, respectively. The i^{th} observation in each simulation is generated as $y_i = (\prod_j x_{ij}) + \mathcal{N}(0,d^2)$ and $y_i = (\sum_j x_{ij}^2) + \mathcal{N}(0,(1.5d)^2)$, respectively, where d is the dimension. For a randomly chosen 0.5% of the training data, the observations are multiplied by a factor of 20. The generalized isotonic models are compared to the

results of a Huber regression, and performance here is measured by mean squared error. Note that we assume that squared error loss represents the true objective performance; the models are fit using Huber loss in order to avoid sensitivity to outliers. The regularized model generated by the minimum loss along the GIRP curve (GIRP Min Huber) is compared with the final GIRP model (GIRP Final Huber) and with the Huber regression model. Table 3 demonstrates that Huber isotonic regression works well with a reasonable number of variables (2-5 for the first simulation and 2-4 for the second simulation), however, again, a simple Huber regression outperforms GIRP for higher dimensions due to overfitting. An important note here is the effect of regularization. The average loss using the unregularized isotonic model is not statistically superior at any dimension to the average loss using a Huber regression while the regularized isotonic model produces statistically improved performance.

Figures 2 and 3 display regularization paths for the Poisson and Huber simulations, respectively. Each curve shows the performance from using increasingly complex models generated by GIRP. Take, for example, the first curve (d=2) under Model 1 in Figure 2. The x-axis states the number of partitions in the particular GIRP model and the y-axis measures the negative Poisson log-likelihood of using the GIRP models (trained on the subtraining data) to predict the validation data. As the number of partitions increases (i.e., as the model becomes more complex), performance improves. Consider next d=5 under the same model. After 12 iterations of GIRP the performance begins to worsen (the minimum along each curve is shown by a diamond). This is exactly the effect of regularization. Performance improves as the model complexity increases up to a certain point at which increasing the complexity further overfits the model and performance declines. Thus, as done to obtain the performance in Tables 2 and 3, the model along the path that gives the best performance on the validation data is used to make predictions on the independent testing data.

The curves in Figure 3 show similar paths for the generalized isotonic regressions with Huber loss where performance is measured by mean squared error. Here the effects of regularization are

much more pronounced than they are in the Poisson simulations. This suggests that robust regressions on applications where isotonicity is desired would greatly benefit from the regularization of GIRP with Huber loss. We next exhibit this robustness effect on a data set for predicting the miles-per-gallon of automobiles.

Model 1: $y_i \sim \text{Poisson}(\prod_i \sqrt{x_{ij}})$ with $x_{ij} \sim \mathcal{U}[0, 10]$

Dim	GIRP Min Poisson	GIRP Final Poisson	Poisson Regression	Min	GIRP
	Neg. Log-Likelihood	Neg. Log-Likelihood	Neg. Log-Likelihood	Path	Length
2	$30678.65 \ (\pm \ 23.83)$	$30678.73~(\pm~23.84)$	32031.44 (± 24.22)	217	298
3	$36395.90 (\pm 38.03)$	$36397.99 \ (\pm\ 37.84)$	40421.91 (± 37.30)	90	618
4	$43908.67 (\pm 54.93)$	$43944.68 \ (\pm \ 56.47)$	$54108.80 (\pm 60.77)$	78	584
5	66812.67 (\pm 240.06)	68096.06 (± 347.41)	$81096.57 (\pm 132.33)$	30	371
6	$200068.80 (\pm 1072.14)$	$220308.20 \ (\pm \ 2059.38)$	$140478.56 (\pm 398.51)$	9	479

Model 2: $y_i \sim \text{Poisson}(\sum_i x_{ij}^2)$ with $x_{ij} \sim \mathcal{U}[5, 10]$

Dim	GIRP Min Poisson	GIRP Final Poisson	Poisson Regression	Min	GIRP
	Neg. Log-Likelihood	Neg. Log-Likelihood	Neg. Log-Likelihood	Path	Length
2	56957.75 (± 31.19)	56957.77 (± 31.20)	57802.85 (± 31.17)	661	794
3	60650.13 (\pm 30.83)	$60650.62~(\pm~31.32)$	$60861.90 (\pm 29.95)$	105	1239
4	$64008.55 (\pm 40.17)$	$64041.56 (\pm 44.10)$	62956.84 (± 23.68)	57	1105
5	$67837.30 (\pm 50.78)$	$68182.67 (\pm 78.18)$	64590.51 (± 30.33)	16	806
6	$74438.33 \ (\pm \ 97.72)$	$75479.73 (\pm 91.07)$	65936.54 (± 28.06)	16	544

Table 2: Statistics for count data simulations generated by two different models as labeled above. *GIRP Min (Final) Poisson Neg. Log-Likelihood* (LL) refers to the negative Poisson LL of predicting on independent testing data using the model that produced the minimum (final) loss along a regularization path generated on training data. *Min Path* is the number of partitions made to generate the minimum negative Poisson LL and GIRP Path Length is the number of partitions in the global generalized isotonic solution. *Poisson Regression Neg. LL* is the negative Poisson LL from using Poisson regressions. Bolded MSE values for minimum and final GIRP negative Poisson LL indicate that they are significantly lower than the negative LL of the Poisson regression at level .05.

3.3 Predicting Miles-Per-Gallon

The next example compares the l_2 versus Huber loss regressions. Note that Huber loss function is an example that cannot be solved using the theory of Barlow & Brunk (1972) and l_2 isotonic regression. This example uses a data set of 392 automobiles (Frank & Asuncion 2010) and models miles-per-gallon using the following seven variables: origin, model year, number of cylinders,

Model 1: $y_i = (\prod_j x_{ij}) + \mathcal{N}(0, d^2)$ with $x_{ij} \sim \mathcal{U}[0, 3]$

Dim	GIRP Min Huber	GIRP Final Huber	Huber Regression	Min	GIRP
	MSE	MSE	MSE	Path	Length
2	$4.21 (\pm 0.30)$	$4.35~(\pm~0.36)$	$4.55 (\pm 0.03)$	49	421
3	9.69 (\pm 0.07)	1 1.71 (± 2.44)	$13.18 (\pm 0.10)$	27	1607
4	22.93 (\pm 0.26)	$90.83 (\pm 64.78)$	$36.94 (\pm 0.51)$	10	3645
5	83.20 (\pm 1.23)	$280.08 (\pm 106.02)$	$115.47 (\pm 2.43)$	6	5783
6	$370.56 (\pm 10.41)$	$2080.71 (\pm 915.59)$	391.01 (± 12.09)	3	7531

Model 2: $y_i = (\sum_i x_{ij}^2) + \mathcal{N}(0, (1.5d)^2)$ with $x_{ij} \sim \mathcal{U}[0, 5]$

	9.		, , , ,	, ,	
Dim	Huber Min Huber	GIRP Final Huber	Huber Regression	Min	GIRP
	MSE	MSE	MSE	Path	Length
2	9.60 (\pm 0.07)	$14.12 (\pm 8.35)$	$15.98 (\pm 0.11)$	57	1154
3	$23.79 \ (\pm \ 0.20)$	$60.17 (\pm 34.96)$	$30.61 (\pm 0.23)$	33	3283
4	48.20 (± 0.41)	193.37 (± 83.58)	$50.02 (\pm 0.36)$	16	5705
5	$85.62 (\pm 0.58)$	599.59 (± 298.06)	$73.44 (\pm 0.54)$	8	7785
6	$145.06 (\pm 1.34)$	$1602.43 (\pm 620.45)$	$101.12 (\pm 0.73)$	8	9283

Table 3: Statistics for count data simulations generated by two different models as labeled above. *GIRP Min (Final) Huber MSE* refers to the MSE of predicting on independent testing data using the model that produced the minimum (final) loss along a regularization path generated on training data. *Min Path* is the number of partitions made to generate the minimum MSE and GIRP Path Length is the number of partitions in the global generalized isotonic solution. *Huber Regression MSE* is the MSE from using Huber regressions.

acceleration, displacement, horsepower, and weight. Isotonic regression with an l_2 loss function was already shown to be useful for this data set in Luss et al. (2012). In this experiment, we have modified one data point to be an outlier. This random point is chosen such that isotonicity constraints require its fit to be less than the fits of five other data points. The experiment simulates a real-life outlier problem which affects the training sample but should not affect prediction. We assume squared error loss to be the true prediction criterion (therefore the out-of-sample evaluation criterion), and fit models with Huber loss to avoid sensitivity to outliers. Table 4 displays the results of one random division of the data (2/3 for training, 1/3 for testing). A paired t-test comparing the out-of-sample predictive performance of the two models (IRP and GIRP) confirms the significant edge of the model generated with Huber's loss function in this setting.

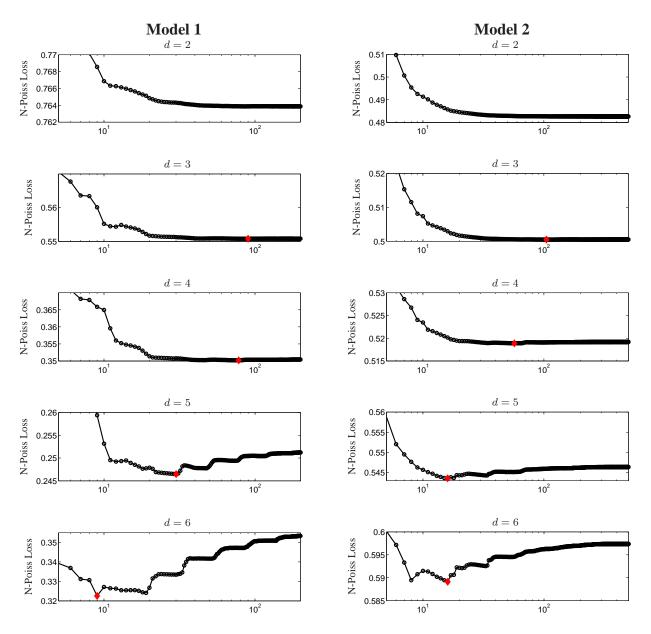


Figure 2: Normalized negative Poisson log-likelihood (LL) for predictions of count data simulations with different dimensions d. Each path is normalized by the loss of the initial model. The x-axis in each figure corresponds to the number of partitions made by GIRP, i.e., the curves show how the normalized negative Poisson LL of test data varies as the GIRP algorithm progresses. Model 1 uses the function $y_i \sim \text{Poisson}(\prod_j \sqrt{x_{ij}})$ with $x_{ij} \sim \mathcal{U}[0,10]$ and Model 2 uses the function $y_i \sim \text{Poisson}(\sum_j x_{ij}^2)$ with $x_{ij} \sim \mathcal{U}[0,5]$. Fifty simulations were run with 12000 training and 3000 testing points. Only the first few hundred partitions of the paths are displayed in order to make the loss of the earlier GIRP iterations visually clearer. Scales also differ in order to make the shapes of the curves clear.

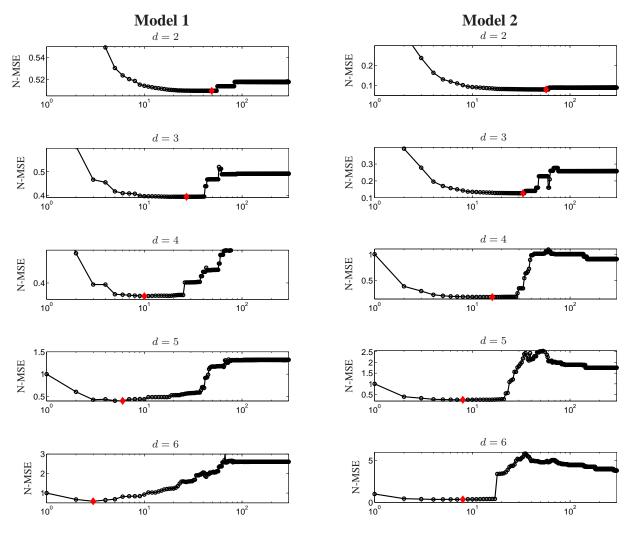


Figure 3: Normalized MSE for predictions of two isotonic simulations with different dimensions d. Each path is normalized by the MSE of the initial model. The x-axis in each figure corresponds to the number of partitions made by GIRP, i.e., the curves show how the normalized MSE of test data varies as the GIRP algorithm progresses. Model 1 uses the function $y_i = (\prod_j x_{ij}) + \mathcal{N}(0, d^2)$ with $x_{ij} \sim \mathcal{U}[0, 10]$ and Model 2 uses the function $y_i = (\sum_j x_{ij}^2) + \mathcal{N}(0, (1.5d)^2)$ with $x_{ij} \sim \mathcal{U}[0, 5]$. Fifty simulations were run with 12000 training and 3000 testing points. Only the first few.

4 Conclusion

In this paper, we show how a relatively minor adjustment to the previously proposed IRP algorithm leads to a generalization allowing us to efficiently fit isotonic models under any convex differentiable loss function. Our proposed GIRP algorithm also generates regularized isotonic solutions along its path, in addition to the optimal isotonic solution. An important remaining challenge is

Number	IRP LS	GIRP Huber	IRP LS	IRP LS	GIRP Huber	GIRP Huber
Variables	Min MSE	Min MSE	Min Path	Path Length	Min Path	Path Length
1	37.47 ± 9.67	38.48 ± 10.42	2	3	2	3
2	31.22 ± 7.04	$\textbf{27.01} \pm \textbf{6.28}$	17	17	7	16
3	21.19 ± 6.75	15.83 ± 4.76	12	30	5	30
4	22.90 ± 6.78	15.53 ± 4.01	4	53	11	54
5	19.94 ± 7.06	10.95 ± 3.15	4	78	29	81
6	17.55 ± 5.99	9.78 ± 2.89	4	86	69	90
7	18.91 ± 6.29	10.24 ± 3.51	4	95	71	95

Table 4: Statistics for auto mpg data. Miles-per-gallon is regressed on a seven potential variables: origin, model year, number of cylinders, acceleration, displacement, horsepower, and weight. A comparison between the results of IRP and GIRP with Huber loss is shown. One data point y_i such that $x_i \leq x_j$ for $j=1\ldots 5$ was modified to be an outlier. δ for Huber loss is set to one standard deviation of the training responses. Bold demonstrates statistical significance with 95% confidence determined by a paired t-test using 131 out-of-sample observed squared losses obtained from models trained on 261 in-sample observations.

to generalize the approach to handling convex non-differentiable loss functions (like absolute loss or the hinge loss of support vector machines), an important topic for future research. Our analysis does not hold in this case due to nonuniqueness of the subproblems.

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5 Appendix

We need the following additional terminology: A group X majorizes (minorizes) another group Y if $X \succeq Y$ ($X \preceq Y$). A group X is a majorant (minorant) of $X \cup A$ where $A = \bigcup_{i=1}^k A_i$ if $X \not\prec A_i$ ($X \not\succ A_i$) $\forall i = 1 \dots k$.

Theorem 1:

Proof. We prove by contradiction. Assume there exists a union of K blocks in V in the optimal solution labeled $\mathcal{M}=M_1\cup\ldots\cup M_K$ that get broken by the cut, with M_1 and M_K as the minorant and majorant block in \mathcal{M} , and M_k^L and M_k^U as the groups in M_k below and above the cut. Define \mathcal{L} as the union of all blocks in V that lie "below" the algorithm cut, \mathcal{U} as the union of all blocks in V that lie "above" the algorithm cut. Further define $A_K^L\subseteq\mathcal{L}$ ($A_1^U\subseteq\mathcal{U}$) as the union of blocks along the algorithm cut such that $A_K^L\succ M_K^L$ ($A_1^U\prec M_1^U$). Figure 4 depicts an example of these definitions where $A_1^U=A_1^L=A_K^U=A_K^L=\{\}$ for simplicity.

We first prove that $w_{M_1} > w_V$. First, consider the case $A_1^U = \{\}$. By convexity of $f_i(\cdot)$ and summing over group M_1^U , we have

$$\sum_{i \in M_1^U} f_i(w_{M_1^U}) \geq \sum_{i \in M_1^U} f_i(w_V) + (w_{M_1^U} - w_V) \sum_{i \in M_1^U} \frac{\partial f_i(\hat{y}_i)}{\partial \hat{y}_i} \bigg|_{w_V}.$$

Definition of the weight operator gives

$$\sum_{i \in M_1^U} f_i(w_{M_1^U}) \le \sum_{i \in M_1^U} f_i(w_V) \Rightarrow (w_{M_1^U} - w_V) \sum_{i \in M_1^U} \frac{\partial f_i(\hat{y}_i)}{\partial \hat{y}_i} \bigg|_{w_V} \le 0.$$

Finally, by the definition of the algorithm cut in (11) since no block exists below M_1^U to affect isotonicity,

$$\sum_{i \in M_i^U} \frac{\partial f_i(\hat{y}_i)}{\partial \hat{y}_i} \bigg|_{w_V} \le 0 \tag{15}$$

so that $w_{M_1^U} \geq w_V$. Since M_1 is a block, we have $w_{M_1^L} > w_{M_1^U}$, and then

$$w_{M_1^L} > w_{M_1^U} > w_V \Rightarrow w_{M_1} > w_V.$$

For the case, $A_1^U \neq \{\}$, we have $w_{M_1} > w_{A_1^U} > w_V$ with the first inequality due to optimality and the second follows directly the proof above replacing M_1^U by A_1^U . A proof for $w_{M_K} < w_V$ follows a similar argument focusing on M_K^L . Putting this together gives $w_{M_1} > w_V > w_{M_K}$, which contradicts that M_1 and M_K are blocks in the global solution, since by assumption then $w_{M_1} < w_{M_K}$. The case K = 1 is also trivially covered by the above arguments. We conclude that the algorithm cannot cut any block.



Figure 4: Illustration of proof of Theorem 1. Black lines separate blocks. The diagonal red line through the center demonstrates a cut of Algorithm 1. \mathcal{L} is the union of blue blocks below the cut and \mathcal{U} is the union of green blocks above the cut. White blocks are blocks that are potentially split by Algorithm 1. These blocks are split into M_1^L,\ldots,M_5^L below the cut and M_1^U,\ldots,M_5^U above the cut. In the proof, $M_i=M_i^L\cup M_i^U$ $\forall i=1\ldots 5$. The proof shows, for example, that if the algorithm splits M_1 into M_1^L and M_1^U according to the defined cut in 11, then there must be no isotonicity violation when creating blocks from M_1^L and M_1^U . However, since M_1 is assumed to be a block, there must exist an isotonicity violation between M_1^L and M_1^U , providing a contradiction.

Theorem 2:

Proof. The proof is by induction. The base case, i.e., first iteration, where all points form one group is trivial. The first cut is made by solving linear program (11) which constrains the solution to maintain isotonicity.

Assuming that iteration k (and all previous iterations) provides an isotonic solution, we prove that iteration k+1 must also maintain isotonicity. Figure 5 helps illustrate the situation described here. Let G be the group split at iteration k+1 and denote A (B) as the group under (over) the cut. Let $A = \{X : X \text{ is a group at iteration } k+1, \exists i \in X \text{ such that } (i,j) \in \mathcal{I} \text{ for some } j \in A\}$ (i.e., $X \in \mathcal{A}$ border A from below).

Consider iteration k + 1. Denote $\mathcal{X} = \{X \in \mathcal{A} : w_A < w_X\}$ (i.e., $X \in \mathcal{X}$ violates isotonicity with A). The split in G causes the fit in nodes in A to decrease. Proof that

$$\left. \sum_{i \in A} \frac{\partial f_i(\hat{y}_i)}{\partial \hat{y}_i} \right|_{w_G} \ge 0$$

follows the proof of (15) in Theorem 1 above so that $w_A \ge w_G$. We will prove that when the fits in A decrease, there can be no groups below A that become violated by the new fits to A, i.e., the decreased fits in A cannot be such that $\mathcal{X} \ne \{\}$.

We first prove that $\mathcal{X} = \{\}$ by contradiction. Assume $\mathcal{X} \neq \{\}$. Denote $k_0 < k+1$ as the iteration at which the last of the groups in \mathcal{X} , denoted D, was split from G and suppose at iteration k_0 , G was part of a larger group H and D was part of a larger group F. It is important to note that $X \cap (F \cup H) = \{\} \ \forall X \in \mathcal{X} \setminus D$ at iteration k_0 because by assumption all groups in $\mathcal{X} \setminus D$ were separated from A before iteration i. Thus, at iteration k_0 , D is the only group bordering A that violates isotonicity.

Let D_U denote the union of D and all groups in F that majorize D. By construction, D_U is a majorant in F. Hence $w_{D_U} < w_{F \cup H}$ by Algorithm 1 and $w_A < w_{D_U}$ by definition since $w_{D_U} > w_D > w_A$. Also by construction, any set $X \in H$ that minorizes A has $w_X < w_A$ (each set X that minorizes A besides D such that $w_X < w_A$ has already been split from A). Hence we can

denote A_L as the union of A and all groups in H that minorize A and we have $w_A > w_{A_L}$ and A_L is a minorant in H. Since $A_L \subseteq H$ at iteration i, we have

$$w_{F \cup H} < w_{A_L} < w_A < w_{D_U} < w_{F \cup H}$$

which is a contradiction, and hence the assumption $\mathcal{X} \neq \{\}$ is false. The first inequality is because the algorithm left A_L in H when F was split from H, and the remaining inequalities are due to the above discussion. Hence the split at iterations k+1 could not have caused a break in isotonicity.

A similar argument can be made to show that the increased fit for nodes in B does not cause any isotonic violation. The proof is hence completed by induction.

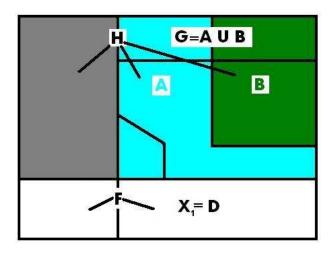


Figure 5: Illustration of proof of Theorem 2 showing the defined sets at iteration k+1. G is the set divided at iteration k+1 into A (all blue area) and B (all green area). The group bordering A from below denoted by X_1 (also referred to as D in the proof) is in violation with A. At iteration k_0 , G is part of the larger group H and X_1 is part of the larger group F. At iteration k_0 , groups F and H are separated. The proof shows that when A and B are split at iteration k+1, no group such as X_1 where $w_{X_1} > w_A$ could have existed. In the picture, X_1 must have been separated at an iteration $k_0 < k+1$, but the proof, through contradiction, shows that this cannot occur.